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Thr Arg Ile Ser Thr Val Pro Ser Gly Ser Leu Ala Leu Asp Ala Ala
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Leu Gly Val Gly Gly Tyr Pro Arg Gly Arg Ile Ile Glu Val Tyr Gly
50 60
Pro Glu Ser Ser Gly Lys Thr Thr Val Ala Leu His Ala Ile Ala Glu
65 70 75 80
Val Gln Gln Gln Gly Gly Gln Ala Ala Phe Ile Asp Ala Glu His Ala
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                 85
Leu Asp Pro Val Tyr Ala Gln Lys Leu Gly Val Asn Ile Asp Glu Leu
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                                 105
                                                       110
Leu Leu Ser Gln Pro Asp Thr Gly Glu Gln Ala Leu Glu Ile Ala Glu
        115
                             120
Ala Leu Val Arg Ser Gly Ala Val Asp Ile Val Val Ile Asp Ser Val
                         135
  130
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Ala Ala Leu Val Pro Lys Ala Glu Ile Glu Gly Asp Met Gly Asp Ser
145 _ _ _ 150 155 _ 160
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His Val Gly Leu Gln Ala Arg Leu Met Ser Gln Ala Leu Arg Lys Leu
165 170 175
                165
Ser Gly Ala Ile Asn Lys Ser Lys Thr Ile Ala Ile Phe Ile Asn Gln
            180
                                 185
Ile Arg Glu Lys Val Gly Val Met Phe Gly Asn Pro Glu Thr Thr Pro
                             200
                                                   205
Gly Gly Arg Ala Leu Lys Phe Tyr Ser Ser Val Arg Leu Glu Val Arg
                         215
                                              22Ō
Arg Ala Glu Gln Leu Lys Gln Gly Asn Asp Val Met Gly Asn Lys Thr
                     230
Lys Ile Lys Val Val Lys Asn Lys Val Ala Pro Pro Phe Arg Thr Ala
                 245
                                      250
                                                           255
Glu Val Asp Ile Met Tyr Gly Glu Gly Ile Ser Lys Glu Gly Glu Ile
260 265 270
lle Asp Leu Gly Thr Glu Leu Asp Ile Val Gln Lys Ser Gly Ala Trp
        275
                             280
                                                  285
Tyr Ser Tyr Gln Glu Glu Arg Leu Gly Gln Gly Arg Glu Asn Ala Lys
   290
                         295
                                              30Ō
Gin Phe Leu Lys Glu Asn Lys Asp Ile Leu Leu Met Ile Gin Glu Gin
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                                          315
Ile Arg Glu His Tyr Gly Leu Asp Thr Gly Gly Ala Ala Pro Ala Gln
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Glu Asp Glu Ala Gln Ala Gln Glu Glu Leu Glu Phe
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Provisional Application 60/535,988 sequence 5502 is identical to SEQ ID NO 6016 of US Application number 10/983,128.